In the Claims

Claims 1-20 (canceled).

Claim 21 (currently amended):

A method of controlling or inhibiting an insect wherein said method comprises contacting said insect with effective amounts of a Protein A, a Protein B, and a Protein C, wherein

- said Protein A is approximately 230-290 kDa, said Protein A consists essentially ofis-a complex forming protein, wherein a polynucleotide A that encodes said Protein A hybridizes under stringent conditions with the full complement of a nucleic acid sequence A that encodes SEQ ID NO:34 (XptA2xwi);
- said Protein B is approximately 130-180 kDa, said Protein B is a complex-forming protein consisting essentially of an, wherein a polynucleotide B that encodes said Protein B hybridizes under stringent conditions with the full complement of a nucleic acid sequence B that encodes a B amino acid sequence selected from the group consisting of SEQ ID NO:22 (TcdB1), SEQ ID NO:45 (TcdB2), and SEQ ID NO:56 (TcaC), SEQ ID NO:18 (XptC1_{Xw}), and SEQ ID NO:49 (XptB1_{Xw});
- said Protein C is approximately 90-120 kDa, said Protein C is a complex-forming protein consisting essentially of an, wherein a polynucleotide C that encodes said Protein C hybridizes under stringent conditions with the full complement of a nucleic acid sequence C that encodes a C amino acid sequence selected from the group consisting of SEQ ID NO:25 (TccC1), SEQ ID NO:47 (TccC3), and SEQ ID NO:57(TccC5), SEQ ID NO:16 (XptB1xwt), and SEQ ID NO:51 (XptC1xwt);
- said Protein A has activity against an insect and said activity is potentiated by said Protein B and said Protein C; and
- said Protein B and said Protein C potentiate the activity of said Protein A:
- wherein when said C amino acid sequence is selected from the group consisting of SEQ ID NO:16 (XptB1_{Xw}) and SEQ ID NO:51 (XptC1_{Xh}), said B amino acid sequence is selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), and SEQ ID NO:56 (TeaC);

wherein when said B amino acid sequence is selected from the group consisting of SEQ ID NO:18 (XptC1_{Xwi}) and SEQ ID NO:49 (XptB1_{Xb}), said C amino acid sequence is selected from the group consisting of SEQ ID NO:25 (TecC1), SEQ ID NO:47 (TecC3), and SEQ ID NO:57(TecC5) and

wherein said stringent conditions are 0.1X SSC and 0.1% SDS at 55° C.

Claim 22 (currently amended):

The method of claim 21 wherein said Protein $\underline{C}A$ -comprises SEQ ID NO: $\underline{47}$ 34 ($\underline{\text{TccC}3XptA2}_{Xwi}$).

Claim 23 (currently amended):

The method of claim 21 wherein said <u>Protein B comprises amino acid sequence is-SEQ ID NO:45 (TcdB2).</u>

Claim 24 (currently amended):

The method of claim 21 wherein said TccC5) and SEQ ID NO:57 (TccC5).

Claim 25 (currently amended):

The method of claim 21 wherein said <u>Protein nucleic acid sequence-B comprises encodes</u> SEQ ID NO:45 (TcdB2), and <u>Protein nucleic acid sequence-C comprises encodes-SEQ ID NO:47 (TccC3).</u>

Claims 26-33 (canceled).

Claim 34 (currently amended):

A method of inhibiting an insect wherein said method comprises contacting said insect with an A component and a B component, and a C component, wherein said components form an insecticidal toxin complex, wherein

- said A component is a 230-290 kDa complex-forming protein having at least 99 95% identity with an A amino acid sequence selected from the group consisting of SEQ ID NO:34 (XptA2)-and-SEQ ID NO:14 (XptA1);
- said B component is a 130-180 kDa complex-forming protein having at least <u>99</u> 95% identity with a B amino acid sequence selected from the group consisting of SEQ ID NO:22 (TcdB1), SEQ ID NO:45 (TcdB2), and SEQ ID NO:56 (TcaC), SEQ ID-NO:18 (XptC1_{xw1}), and SEQ ID-NO:49 (XptB1_{xw2});
- said C component is a 90-120 kDa complex-forming protein having at least 95% identity with a C amino acid sequence selected from the group-consisting of SEQ ID NO:25 (TeeC1), SEQ ID NO:47 (TeeC3), SEQ ID NO:57(TeeC5), SEQ ID NO:16 (XptB1xw); and SEQ ID NO:51 (XptC1xb);
- wherein said A component has activity against an insect, and wherein said B component is a potentiator of said A component and C components potentiate said activity:
- wherein when said C amino acid sequence is selected from the group consisting of SEQ ID NO:16 (XptB1_{Xwi}) and SEQ ID NO:51 (XptC1_{Xb}), said B amino acid sequence is selected from the group consisting of SEQ ID NO:22 (TcdB1), SEQ ID NO:45 (TcdB2), and SEQ ID NO:56 (TcaC); and
- wherein when said B amino acid sequence is selected from the group consisting of SEQ ID NO:18 (XptC1_{Xw}) and SEQ ID NO:49 (XptB1_{Xb})_x said C amino acid sequence is selected from the group consisting of SEQ ID NO:25 (TeeC1), SEQ ID NO:47 (TeeC3), and SEQ ID NO:57 (TeeC5).

Claim 35 (currently amended):

The method of claim 34 wherein said A $\underline{\text{component amino acid-sequence}}$ is SEQ ID NO:34 (XptA2).

Claim 36 (currently amended):

A method of inhibiting an insect wherein said method comprises contacting said insect with an A component, a B component, and a C component, wherein said components form an insecticidal toxin complex, wherein

- said A component is a 230-290 kDa complex-forming protein having at least 95% identity with SEQ ID NO:34 (XptA2)an A sequence selected from the group consisting of SEQ ID NO:21 (TedA), SEQ ID NO:62 (TedA2), SEQ ID NO:63 (TedA4), and SEO ID NO:59 (TebA1);
- said B component is a 130-180 kDa complex forming protein having at least 95% identity with an amino acid sequence selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), SEQ ID NO:56 (TeaC), SEQ ID NO:18 (XptC1xw), and SEQ ID NO:49 (XptB1xb);
- said C component is a 90-120 kDa complex-forming protein having at least 95% identity with an amino acid sequence selected from the group consisting of SEQ ID NO:25 (TccC1), SEQ ID NO:47 (TccC3), and SEQ ID NO:57 (TccC5), SEQ ID NO:16 (XptB1xw,), and SEQ ID NO:51 (XptC1xb);
- wherein said A component has activity against an insect, and said B and C component is

 a potentiator of said A component, and any differences between said A component and SEQ ID NO:34, and between said C component and said amino acid sequence, are conservative amino acid substitutionseomponents potentiate said toxin activity;
- wherein when said C sequence is selected from the group consisting of SEQ ID NO:25

 (TecC1), SEQ ID NO:47 (TecC3), and SEQ ID NO:57 (TecC5), said B sequence
 is selected from the group consisting of SEQ ID NO:18 (XptC1_{Xw}) and SEQ ID
 NO:49 (XptB1_{Xh}) when said C sequence is selected from the group consisting of
 SEQ ID NO:25 (TecC1), SEQ ID NO:47 (TecC3), and SEQ ID NO:57 (TecC5);
 and
- wherein when said B sequence is selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), and SEQ ID NO:56 (TeaC), said C sequence is

selected from the group consisting of SEQ ID NO:16 (XptB1_{Xm}) and SEQ ID NO:51 (XptC1_{Xb}) when said B sequence is selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), and SEQ ID NO:56 (TeaC).

Claim 37 (currently amended):

The method of claim 36 wherein said \underline{C} component comprises A sequence is SEQ ID NO:47 $\underline{24}$ (TccC3 \underline{T} edA).

Claim 38 (currently amended):

The method of claim 34, wherein said method further comprises contacting said insect with a

- said A component comprises an amino acid sequence selected from the group consisting of SEQ ID-NO:34 (XptA2) and SEQ ID-NO:14 (XptA1);
- said B component comprises an amino acid sequence selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), SEQ ID NO:56 (TeaC), SEQ ID NO:18 (XptC1_{Xwi}), and SEQ ID NO:49 (XptB1_{Xb}); and
- said C component <u>comprising emprises</u>-an amino acid sequence selected from the group consisting of SEQ_ID_NO:25 (TeeC1), SEQ_ID_NO:47 (TeeC3), SEQ_ID_NO:57(TeeC5), SEQ_ID_NO:16 (XptB1_{Xwi}), and SEQ_ID_NO:51 (XptC1_{Xb});
- wherein when said C component <u>comprises an amino acid sequence selected from the group consisting of SEQ ID NO:16 (XptB1_{Xwt}) and SEQ ID NO:51 (XptC1_{Xb}); said B component comprises an amino acid sequence selected from the group consisting of SEQ ID NO:22 (TedB1), SEQ ID NO:45 (TedB2), and SEQ ID NO:56 (TeaC); and</u>
- wherein when said B component is selected from the group consisting of SEQ ID NO:18

 (XptC1_{xwi}) and SEQ ID NO:49 (XptB1_{xki}), said C component comprises an amino acid sequence selected from the group consisting of SEQ ID NO:25 (TeeC1), SEQ ID NO:47 (TeeC3), and SEQ ID NO:57(TeeC5).

Claim 39 (currently amended):

The method of claim 38 36 wherein when said method further comprises contacting said insect with a B.A-component selected from the group consisting of SEQ ID NO:18 (XptC1_{Xxi} and SEQ ID NO:49 (XptB1_{Xx})comprises SEQ-ID NO:34 (XptA2).

Claim 40 (currently amended):

The method of claim 35 wherein said B <u>component amino acid sequence</u> is SEQ ID NO:45 (TcdB2) and said C amino acid sequence is selected from the group consisting of SEQ ID NO:47 (TeeC3) and SEO ID NO:57 (TeeC5).

Claim 41 (currently amended):

The method of claim $\underline{36}$ 40 wherein said \underline{A} component C amino acid-sequence is SEQ ID NO:34.7 (XptA2 $\overline{\ }$ 4ceC3).

Claim 42 (currently amended):

The method of claim 39 wherein said <u>C-B</u> component comprises SEQ ID NO:45 (TcdB2), and said C component comprises SEQ ID NO:47 (TccC3).